SEQUENCE LISTING

<110> Paszty, Christopher
Gao, Yongming

<120> Cystine Knot Polypeptides: Cloaked-2 Molecules and Uses Thereof

<130> 01017/37428A

<140> US 10/679,670
<141> 2003-10-06

<150> US 60/208,550
<151> 2000-06-01

<150> US 60/223,542
<151> 2000-08-04

<160> 25

<170> PatentIn version 3.0

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<212> DNA
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ttcaagaatg atgccacgga aatcatcccc gagctcggag agtaccccga gcctccaccg 180

gagctggaga acaacaagac catgaaccgg gcggagaacg gagggcggcc tccccaccac 240

ccctttgaga ccaaagacgt gtccgagtac agctgccgcg agctgcactt cacccgctac 300

gtgaccgatg ggccgtgccg cagcgccaag ccggtcaccg agctggtgtg ctccggccag 360

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cccggtggtg aggcgccgcg cgcgcgcaag gtgcgcctgg tggcctcgtg caagtgcaag 540

cgcctcaccc gcttccacaa ccagtcggag ctcaaggact tcgggaccga ggccgctcgg 600

ccgcagaagg gccggaagcc gcggccccgc gcccggagcg ccaaagccaa ccaggccgag 660

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<213> Homo sapiens

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Leu Gly Glu Tyr Pro Glu Pro Pro Pro Glu Leu Glu Asn Asn Lys Thr
20 25 30

Met Asn Arg Ala Glu Asn Gly Gly Arg Pro Pro His His Pro Phe Glu
35 40 45

Thr Lys Asp Val Ser Glu Tyr Ser Cys Arg Glu Leu His Phe Thr Arg
50 55 60

Tyr Val Thr Asp Gly Pro Cys Arg Ser Ala Lys Pro Val Thr Glu Leu
65 70 75 80

Val Cys Ser Gly Gln Cys Gly Pro Ala Arg Leu Leu Pro Asn Ala Ile
85 90 95

Gly Arg Gly Lys Trp Trp Arg Pro Ser Gly Pro Asp Phe Arg Cys Ile
100 105 110

Pro Asp Arg Tyr Arg Ala Gln Arg Val Gln Leu Leu Cys Pro Gly Gly
115 120 125

Glu Ala Pro Arg Ala Arg Lys Val Arg Leu Val Ala Ser Cys Lys Cys
130 135 140

Lys Arg Leu Thr Arg Phe His Asn Gln Ser Glu Leu Lys Asp Phe Gly
145 150 155 160

Thr Glu Ala Ala Arg Pro Gln Lys Gly Arg Lys Pro Arg Pro Arg Ala
165 170 175

Arg Ser Ala Lys Ala Asn Gln Ala Glu Leu Glu Asn Ala Tyr
180 185 190

<210> 3
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<212> DNA
<213> Mus musculus

<400> 3

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ggagagtacc ccgagcctcc tcctgagaac aaccagacca tgaaccgggc ggagaatgga 180

ggcagacctc cccaccatcc ctatgacgcc aaagatgtgt ccgagtacag ctgccgcgag 240

ctgcactaca cccgcttcct gacagacggc ccatgccgca gcgccaagcc ggtcaccgag 300

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aagtggtggc gcccgaacgg accggatttc cgctgcatcc cggatcgcta ccgcgcgcag 420

cgggtgcagc tgctgtgccc cgggggcgcg gcgccgcgct cgcgcaaggt gcgtctggtg 480

gcctcgtgca agtgcaagcg cctcacccgc ttccacaacc agtcggagct caaggacttc 540

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aaagccaacc aggcggagct ggagaacgcc tactag 636

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Leu Gly Glu Tyr Pro Glu Pro Pro Pro Glu Asn Asn Gln Thr Met Asn
20 25 30

Arg Ala Glu Asn Gly Gly Arg Pro Pro His His Pro Tyr Asp Ala Lys
35 40 45

Asp Val Ser Glu Tyr Ser Cys Arg Glu Leu His Tyr Thr Arg Phe Leu
50 55 60

Thr Asp Gly Pro Cys Arg Ser Ala Lys Pro Val Thr Glu Leu Val Cys
65 70 75 80

Ser Gly Gln Cys Gly Pro Ala Arg Leu Leu Pro Asn Ala Ile Gly Arg
85 90 95

Val Lys Trp Trp Arg Pro Asn Gly Pro Asp Phe Arg Cys Ile Pro Asp
100 105 110

Arg Tyr Arg Ala Gln Arg Val Gln Leu Leu Cys Pro Gly Gly Ala Ala
115 120 125

Pro Arg Ser Arg Lys Val Arg Leu Val Ala Ser Cys Lys Cys Lys Arg
130 135 140

Leu Thr Arg Phe His Asn Gln Ser Glu Leu Lys Asp Phe Gly Pro Glu
145 150 155 160

Thr Ala Arg Pro Gln Lys Gly Arg Lys Pro Arg Pro Gly Ala Lys Ala
165 170 175

Asn Gln Ala Glu Leu Glu Asn Ala Tyr
180 185

<210> 5
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<213> Homo sapiens

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Met Gln Leu Pro Leu Ala Leu Cys Leu Val Cys Leu Leu Val His Thr
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20 25 30

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Ala Thr Glu Ile Ile Pro Glu Leu Gly Glu Tyr Pro Glu Pro Pro Pro
        35                  40                  45

Glu Leu Glu Asn Asn Lys Thr Met Asn Arg Ala Glu Asn Gly Gly Arg
    50                  55                  60

Pro Pro His His Pro Phe Glu Thr Lys Asp Val Ser Glu Tyr Ser Cys
65                  70                  75                  80

Arg Glu Leu His Phe Thr Arg Tyr Val Thr Asp Gly Pro Cys Arg Ser
                85                  90                  95

Ala Lys Pro Val Thr Glu Leu Val Cys Ser Gly Gln Cys Gly Pro Ala
            100                 105                 110

Arg Leu Leu Pro Asn Ala Ile Gly Arg Gly Lys Trp Trp Arg Pro Ser
        115                 120                 125

Gly Pro Asp Phe Arg Cys Ile Pro Asp Arg Tyr Arg Ala Gln Arg Val
    130                 135                 140

Gln Leu Leu Cys Pro Gly Gly Glu Ala Pro Arg Ala Arg Lys Val Arg
145                 150                 155                 160

Leu Val Ala Ser Cys Lys Cys Lys Arg Leu Thr Arg Phe His Asn Gln
                165                 170                 175

Ser Glu Leu Lys Asp Phe Gly Thr Glu Ala Ala Arg Pro Gln Lys Gly
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Arg Lys Pro Arg Pro Arg Ala Arg Ser Ala Lys Ala Asn Gln Ala Glu
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Leu Glu Asn Ala Tyr
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<400> 6

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Ala Phe Cys Ala Val Glu Gly Gln Gly Trp Gln Ala Phe Arg Asn Asp
            20                  25                  30

Ala Thr Glu Val Ile Pro Gly Leu Gly Glu Tyr Pro Glu Pro Pro Pro
        35                  40                  45

Glu Asn Asn Gln Thr Met Asn Arg Ala Glu Asn Gly Gly Arg Pro Pro
    50                  55                  60

His His Pro Tyr Asp Ala Lys Asp Val Ser Glu Tyr Ser Cys Arg Glu
65                  70                  75                  80

Leu His Tyr Thr Arg Phe Leu Thr Asp Gly Pro Cys Arg Ser Ala Lys
                85                  90                  95

Pro Val Thr Glu Leu Val Cys Ser Gly Gln Cys Gly Pro Ala Arg Leu
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            100                 105                 110

Leu Pro Asn Ala Ile Gly Arg Val Lys Trp Trp Arg Pro Asn Gly Pro
        115                 120                 125

Asp Phe Arg Cys Ile Pro Asp Arg Tyr Arg Ala Gln Arg Val Gln Leu
    130                 135                 140

Leu Cys Pro Gly Gly Ala Ala Pro Arg Ser Arg Lys Val Arg Leu Val
145                 150                 155                 160

Ala Ser Cys Lys Cys Lys Arg Leu Thr Arg Phe His Asn Gln Ser Glu
                165                 170                 175

Leu Lys Asp Phe Gly Pro Glu Thr Ala Arg Pro Gln Lys Gly Arg Lys
            180                 185                 190

Pro Arg Pro Gly Ala Lys Ala Asn Gln Ala Glu Leu Glu Asn Ala Tyr
        195                 200                 205
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<213> Artificial

<220>
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<210> 8
<211> 26
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<213> Artificial sequence

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<210> 9
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<212> DNA
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<210> 10
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cgatccggga tgcagcggaa gtcg 24

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ccatcctaat acgactcact atagggc 27

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ccatcctaat acgactcact atagggc 27

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ggtcaccgag ttggtgtgct c 21

<210> 18
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<400> 18

actcactata gggctcgagc ggc 23

<210> 19
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<400> 20

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<210> 21
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tgtgtctcgt ctgcctgctg gtacaca 27

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<220>
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<400> 22

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<210> 23
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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
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<212> PRT
<213> Artificial sequence

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<213> Homo sapiens

<400> 25

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Val Pro Ala His Pro Ser Ser Asn Ser Thr Leu Asn Gln Ala Arg Asn
            20                  25                  30

Gly Gly Arg His Phe Ser Asn Thr Gly Leu Asp Arg Asn Thr Arg Val
        35                  40                  45

Gln Val Gly Cys Arg Glu Leu Arg Ser Thr Lys Tyr Ile Ser Asp Gly
    50                  55                  60

Gln Cys Thr Ser Ile Ser Pro Leu Lys Glu Leu Val Cys Ala Gly Glu
65                  70                  75                  80

Cys Leu Pro Leu Pro Val Leu Pro Asn Trp Ile Gly Gly Gly Tyr Gly
                85                  90                  95

Thr Lys Tyr Trp Ser Arg Arg Ser Ser Gln Glu Trp Arg Cys Val Asn
            100                 105                 110

Asp Lys Thr Arg Thr Gln Arg Ile Gln Leu Gln Cys Gln Asp Gly Ser
        115                 120                 125

Thr Arg Thr Tyr Lys Ile Thr Val Val Thr Ala Cys Lys Cys Lys Arg
    130                 135                 140

Tyr Thr Arg Gln His Asn Glu Ser Ser His Asn Phe Glu Ser Met Ser
145                 150                 155                 160

Pro Ala Lys Pro Val Gln His His Arg Glu Arg Lys Arg Ala Ser Lys
                165                 170                 175

Ser Ser Lys His Ser Met Ser
            180
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